

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: August 8, 2002, 22:02:08 ; Search time: 414.95 Seconds

(without alignments)
10583.793 Million cell updates/sec

Title: PCT-US01-46559-1

Perfect score: 224

Sequence: 1 ctgcgtccctccatctgg.....tttggaaaaaaaaaaaaaa 2254

Scoring table: IDENTITY_NUC

GAPOP 10.0 , Gapext 1.0

Searched: 1797656 seqs, 10463268293 residues

Total number of hits satisfying chosen parameters: 3595312

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

GenEmbl:*

1: gb_db:*

2: gb_ntg:*

3: gb_in:*

4: gb_cm:*

5: gb_ov:*

6: gb_pat:*

7: gb_ph:*

8: gb_pl:*

9: gb_pr:*

10: gb_ro:*

11: gb_sts:*

12: gb_sy:*

13: gb_un:*

14: gb_vl:*

15: em_ba:*

16: em_fun:*

17: em_hum:*

18: em_in:*

19: em_mu:*

20: em_om:*

21: em_or:*

22: em_ov:*

23: em_pat:*

24: em_ph:*

25: em_pl:*

26: em_ro:*

27: em_sts:*

28: em_vl:*

29: em_htg_hum:*

30: em_htg_inv:*

31: em_htg_other:*

32: em_htgo_inv:*

33: em_htgo_inv:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No. Score Query Length DB ID Description

1	1652.4	73.3	1880	6	AX017982	Sequence	
2	1330	59.0	1330	6	AX017833	Sequence	
3	1160	54	1162	9	BC000848	Homo sapi	
4	1115	4.4	49.5	2523	AK053388	Homo sapi	
5	969	4.4	43.0	992	BC009386	Homo sapi	
6	880	8	39.1	885	9	HSA23666	
7	772	34.3	772	9	BC015316	Homo sapi	
8	557	24.7	2939	9	BC017708	Homo sapi	
9	488	4	21.7	64759	9	AL592078	Human DNA
10	461	2	20.5	2993	9	AK000617	Homo sapi
c	11	422.6	18.7	5281	9	U08191	Human R kap
c	12	406	18.0	61087	2	AC103392	Mus muscu
c	13	371.4	16.5	585	9	AF397158	Homo sapi
c	14	371.4	16.5	2235	9	HUMRGEA	
c	15	371.4	16.5	25606	2	AC010554	Homo sapi
c	16	371.4	16.5	42999	9	HSU13369	
c	17	371.4	16.5	43962	2	AC068192	Homo sapi
c	18	371.4	16.5	61490	2	AC026330	Homo sapi
c	19	371.4	16.5	101641	2	AL15197	Homo sapi
c	20	371.4	16.5	12833	2	AC068881	Homo sapi
c	21	371.4	16.5	136630	2	AC025572	Homo sapi
c	22	371.4	16.5	140653	2	AL355134	Homo sapi
c	23	371.4	16.5	150286	2	AL353644	Homo sapi
c	24	369.8	16.4	18659	6	AX205305	Sequence
c	25	369.8	16.4	18659	6	AX207387	Sequence
c	26	369.8	16.4	18659	9	HSRRN18S	
c	27	368.8	16.4	50872	2	AC084140	Homo sapi
c	28	368.2	16.3	16008	9	M1008	Human 18S r
c	29	368.2	16.3	55939	2	AC091017	Homo sapi
c	30	367	16.3	61490	2	AC026330	Homo sapi
c	31	363.4	16.1	1824	4	AJ311674	Dasypus n
c	32	363.4	16.1	1824	4	EQUUS CAB	
c	33	360.6	16.0	183292	2	AC094121	Rattus no
c	34	355.6	15.8	1825	4	AJ311675	Erinaceus
c	35	345	15.3	21908	2	AC091632	Homo sapi
c	36	343.4	15.2	2062	10	X56744	M.musculus
c	37	342.4	15.2	39133	2	AC105701	Rattus no
c	38	342.4	15.2	214755	2	AC091339	Rattus no
c	39	338.6	15.0	918	10	MURSB51	
c	40	333	14.8	1920	10	RATRGEA	
c	41	331.4	14.7	18659	6	AX025304	Sequence
c	42	331.4	14.7	18659	6	AX207386	Sequence
c	43	331.4	14.7	18659	10	MMRRA18	
c	44	331.4	14.7	22118	10	X82664	M.musculus
c	45	330.4	14.7	1874	10	RNRNNA06	

ALIGNMENTS

RESULT	1	AX017982	1880 bp	DNA	linear	PAT 07-SEP-2000
LOCUS		AX017982	Sequence	217	from Patent	WO9946375:
DEFINITION		AX017982	Accession			
VERSION		AX017982.1	Version			
KEYWORDS						
SOURCE						
ORGANISM						
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;						
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.						
REFERENCE						
AUTHORS						
Pilarsky, C.						
Human nucleic acid sequences from prostate tissue						
Patent: WO 9946375-A 217 16-SEP-1999;						
SCHMITT ARMIN (DE); SPECHT THOMAS (DE); DAHL EDGAR (DE); HINZMANN BERNRD (DE); ROSENTHAL ANDRE (DE); METAGEN GES FUER GENOMFORSCHUNG (DE); PILARSKY CHRISTIAN (DE)						
Location/Qualifiers						
1. 1880						
/organism="Homo sapiens"						
/db_xref="taxon:9606".						

SUMMARIES

Result No.	Score	Query Length	DB ID	Description
1	1.880			

